

GenCorp version 4.5  
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# OM protein - protein search, using sw model

Run on: June 26, 2001, 16:30:29 ; Search time 57.86 Seconds  
(without alignments)  
209,554 Million cell updates/sec

Title: US-09-484-629a-2

Sequence: 1 MRLALRLAARPGGPPTL.....LDSPSYNAWTKGVVFN 200

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 412676 seqs, 60623988 residues

Total number of hits satisfying chosen parameters: 412676

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%

Listing first 45 summaries

## Database :

1: /SIDSR/gcgdata/geneseq/geneseq/AA1980.DAT:\*  
2: /SIDSR/gcgdata/geneseq/geneseq/AA1981.DAT:\*  
3: /SIDSR/gcgdata/geneseq/geneseq/AA1982.DAT:\*  
4: /SIDSR/gcgdata/geneseq/geneseq/AA1983.DAT:\*  
5: /SIDSR/gcgdata/geneseq/geneseq/AA1984.DAT:\*  
6: /SIDSR/gcgdata/geneseq/geneseq/AA1985.DAT:\*  
7: /SIDSR/gcgdata/geneseq/geneseq/AA1986.DAT:\*  
8: /SIDSR/gcgdata/geneseq/geneseq/AA1987.DAT:\*  
9: /SIDSR/gcgdata/geneseq/geneseq/AA1988.DAT:\*  
10: /SIDSR/gcgdata/geneseq/geneseq/AA1989.DAT:\*  
11: /SIDSR/gcgdata/geneseq/geneseq/AA1990.DAT:\*  
12: /SIDSR/gcgdata/geneseq/geneseq/AA1991.DAT:\*  
13: /SIDSR/gcgdata/geneseq/geneseq/AA1992.DAT:\*  
14: /SIDSR/gcgdata/geneseq/geneseq/AA1993.DAT:\*  
15: /SIDSR/gcgdata/geneseq/geneseq/AA1994.DAT:\*  
16: /SIDSR/gcgdata/geneseq/geneseq/AA1995.DAT:\*  
17: /SIDSR/gcgdata/geneseq/geneseq/AA1996.DAT:\*  
18: /SIDSR/gcgdata/geneseq/geneseq/AA1997.DAT:\*  
19: /SIDSR/gcgdata/geneseq/geneseq/AA1998.DAT:\*  
20: /SIDSR/gcgdata/geneseq/geneseq/AA1999.DAT:\*  
21: /SIDSR/gcgdata/geneseq/geneseq/AA2000.DAT:\*  
22: /SIDSR/gcgdata/geneseq/geneseq/AA2001.DAT:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1004	100.0	200	21	AA199370
2	904	90.0	200	21	AA199372
3	687	68.4	205	21	AA199371
4	135	13.4	41	21	AA199373
5	135	13.4	41	21	AA199375
6	126	12.5	1299	21	AA199373
7	125	12.5	176	21	AA199375
8	125	12.5	176	21	AA199375
9	125	12.5	176	21	AA199375
10	125	12.5	176	21	AA199375
11	125	12.5	195	21	AA199375

12	125	12.5	195	21	AA199370
13	125	12.5	266	21	AA199370
14	125	12.5	266	21	AA199370
15	125	12.5	266	21	AA199370
16	123.5	12.3	1643	21	AA199370
17	123.5	12.3	2110	21	AA199370
18	122	12.2	631	21	AA199370
19	121.5	12.1	1888	20	AA199370
20	120.5	12.0	462	11	AA199370
21	118.5	11.8	583	19	AA199370
22	117.5	11.7	311	19	AA199370
23	117.5	11.7	452	19	AA199370
24	117.5	11.7	467	19	AA199370
25	117.5	11.7	493	19	AA199370
26	116.5	11.6	2819	22	AA199370
27	114.5	11.4	186	18	AA199370
28	114.5	11.4	186	20	AA199370
29	114	11.4	314	20	AA199370
30	114	11.4	316	13	AA199370
31	114	11.4	493	13	AA199370
32	113	11.3	1090	21	AA199370
33	112.5	11.2	2442	21	AA199370
34	112	11.2	303	21	AA199370
35	111	11.1	756	21	AA199370
36	111	11.1	1239	20	AA199370
37	110.5	11.0	315	22	AA199370
38	110.5	11.0	318	13	AA199370
39	109	10.9	914	18	AA199370
40	109	10.9	914	21	AA199370
41	109	10.9	1411	17	AA199370
42	108	10.8	251	21	AA199370
43	108	10.8	2096	21	AA199370
44	107.5	10.7	2115	21	AA199370
45	107.5	10.7	2192	18	AA199370

## ALIGNMENTS

RESULT 1  
AA199370  
ID AA199370 standard; Protein; 200 AA.  
AC AA199370;  
DT 19-JUN-2000 (first entry)  
XX  
DE A 5'-OT EST (oxytocin expressed sequence tag) protein.  
KW Oxytocin expressed sequence tag; 5'-OT EST; obesity; fertility; male;  
KW transgenic animal; human late onset obesity; late onset visceral obesity;  
KW male infertility; wasting; anorexia; cachexia; malabsorptive state;  
KW catabolic state; inflammatory condition; Crohn's disease; AIDS wasting;  
KW burn; cancer; bone disease.  
XX  
OS Rattus sp.  
XX  
PN W0200009686-A1.  
XX  
PD 24-FEB-2000.  
XX  
PF 12-AUG-1999; 99MO-GH02658.  
XX  
PR 12-AUG-1998; 98GB-0017566.  
XX  
PR 06-MAY-1999; 99GB-0010522.  
XX  
PA (MED-) MEDICAL RES COUNCIL.  
XX  
PI Robinson ICAF, Stoye JP, Flavell D, Wells SE, Le Tissier P;  
XX WPI: 2000-224331/19.  
XX N-PSDB: AA261510.  
XX

Arabidopsis thaliana  
Arabidopsis thaliana  
Arabidopsis thaliana  
Arabidopsis thaliana  
Arabidopsis thaliana  
Virus gene pro  
Virus gene pro  
JEST, a human SWAP  
A human trichonayal  
Portion of peptide  
Trypanosoma lympho  
Trypanosoma lympho  
Trypanosoma lympho  
Trypanosoma lympho  
Trypanosoma lympho  
Human 07G27 gene  
Trypanosoma cruzi  
Trypanosoma cruzi  
Human stomach carc  
P. falciparum LSA-R  
P. falciparum LSA g  
Human huntingtin-1  
Human cytoskeletal  
Human pancreatic c  
Mouse huntingtin-1  
Human ZCL protein.  
Human secreted pro  
P. falciparum LSA N  
Huntingtin interac  
Human huntingtin-1  
Nucleolar/endosome  
Human OREX ORE2740  
Human OREX ORE1356  
Human NuMA protein  
LexA/NuMA fusion p

PT New anti-obesity polypeptide useful for treating obesity or infertility

PS Claim 1; Page 116-117; 162pp; English.

XX  
XX  
CC The present sequence represents a 5'-OT-EST (oxytocin expressed sequence tag) polypeptide. The 5'-OT EST gene is involved in the control of CC obesity and fertility in males. 5'-OT EST nucleic acids are useful CC for producing transgenic animals. The transgenic animals created serve CC as a model for human late onset obesity and other related disorders and CC are also used for identifying the genetic cause of obesity. Compounds CC which modulate 5'-OT EST expression or activity are useful in the CC treatment or modulation of late onset visceral obesity or male CC infertility particularly in the disorders related to these conditions CC such as wasting, or anorexia, or cachexia associated with prolonged CC illness, or malabsorptive states or catabolic states associated with CC other diseases such as inflammatory conditions, Crohn's disease or CC AIDS wasting, or burns, or cancer, or bone disease.

XX Sequence 200 AA;

SQ

Query Match 100.0%; Score 1004; DB 21; Length 200;  
Best Local Similarity 100.0%; Pred. No. 1.5e-87;  
Matches 200; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 MIRALNRLAARPGOPPLLLIPVGRKTRHDPKSKSVGRYKMPAYDPAELFYLTRY 60  
Db 1 mIraInrLlaarPgOpPlllIpvgRkTrhdPkSkSvgrYkMpaYdPaElfYlTrry 60

OY 61 RGYRETVRALREFLEVRGKLHEAPACVLAEKKOEAIRHEOETAMNNREBNRLOELR 120  
Db 61 rgyreTvralrefleVrgklheArpAcvlAeKKoEaIrHeoEtAmnnRebnrLoElR 120

OY 121 IARLOLEAOBOLROAEVOAOROEQAOWVIOKEDEVILKLOEAKNFITRENLEARIEEA 180  
Db 121 IaRlOlEaObOlRoAeVoAoRoEqaOwViOkEdEvIlKlOeAkNfiTrenLeaRIea 180

OY 181 LDSPKSYMMAAVTKEGOVVVN 200  
Db 181 ldsPkSyMmAvtKeGoVvn 200

RESULT 2 7  
ID AAY69372  
XN AAY69372 standard; Protein; 200 AA.  
XX  
XX AAY69372;  
XX  
DT 19-JUN-2000 (first entry)  
XX  
DE A mouse 5'-OT EST (oxytocin expressed sequence tag) protein.  
XX  
KM Oxytocin expressed sequence tag; 5'-OT EST; obesity; fertility; male;  
KW transgenic animal; human late onset obesity; late onset visceral obesity;  
KM male infertility; wasting; anorexia; cachexia; malabsorptive state;  
KW catabolic state; inflammatory condition; Crohn's disease; AIDS wasting;  
burn; cancer; bone disease.  
XX  
OS Mus sp.  
XX  
PC WO200009686-A1.  
XX  
PD 24-FEB-2000.  
XX  
PF 12-AUG-1999; 99WO-GB02658.  
XX  
PR 12-AUG-1998; 98GB-0017566.  
XX  
PR 06-MAY-1999; 99GB-0010522.  
XX  
PA (MEDICAL) MEDICAL RES COUNCIL.  
PI Robinson ICAT, Stoye JP, Flavell D, Wells SE, Le Tissier P;

Query Match	Best Local Similarity	Score	DB	Length	Matches	Conservative	Mismatches	Indels	Gaps
1	MLRANRLAARGGGPPPTLLPVRGRKRHPDPAPKSKYGRVMPPAVPAELFVLTERRY	90.0%	60	60	179	10	11	0	0
1	mhrainrlaqrgdrrpprlllpvrgkrkhrdpapkskgrvqtpavpaefvltety	89.5%	60	60	179	10	11	0	0
61	ROYRETVRLRRPFTLEVGKLEHARAGVLAERKQAEIRHOELMANNREERRRLQELR	90.0%	120	120	179	10	11	0	0
61	gyrretvrlrlrftldvrkklhearagvlaerkaqatthelmannrderrmqelr	89.5%	120	120	179	10	11	0	0
121	IKRLQEAQAOELROREVOAORAOEQAWOLKEQVLLKQEPKNTFRRENLEARIEEA	90.0%	180	180	179	10	11	0	0
121	ltrlqldeaqagvqkaeqgrqraeeqawvqlkegevlqldeaknfltrrenleareea	89.5%	180	180	179	10	11	0	0
181	LDSPKSYNNAVTKEGOVRN	90.0%	200	200	179	10	11	0	0
181	ldspksynnavtckegqvvrn	89.5%	200	200	179	10	11	0	0
3	AAAY69371	90.0%	205	205	179	10	11	0	0
AAAY69371	standard; Protein: 205 AA.	89.5%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
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19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	0
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19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	90.0%	205	205	179	10	11	0	0
19-JUN-2000	(first entry)	89.5%	205	205	179	10	11	0	



PA (MEDI-) MEDICAL RES COUNCIL.  
XX  
XX Robinson ICAF, Stoye JP, Flavell D, Wells SE, Le Tissier P;  
XX WPI: 2000-224331/19.  
XX  
XX New anti-obesity polypeptide useful for treating obesity or infertility  
XX in mammals -  
XX  
XX Claim 7; Page 94: 162pp; English.  
XX  
XX The present sequence represents a fragment of a mutant 5'-OR-EST (oxytocin  
CC expressed sequence tag) protein. The 5'-OR EST gene is involved in the  
CC control of obesity and fertility in males. 5'-OR EST nucleic acids are  
CC useful for producing transgenic animals. The transgenic animals created  
CC serve as a model for human late onset obesity and other related  
CC disorders and are also used for identifying the genetic cause of obesity.  
CC Compounds which modulate 5'-OR EST expression or activity are useful in  
CC the treatment or modulation of late onset visceral obesity or male  
CC infertility particularly in the disorders related to these conditions  
CC such as wasting, or anorexia, or cachexia associated with prolonged  
CC illness, or malabsorptive states or catabolic states associated with  
CC other diseases such as inflammatory conditions, Crohn's disease or AIDS  
CC wasting, or burns, or cancer, or bone disease.  
XX  
XX Sequence 41 AA:  
SQ  
Query Match 13.4%; Score 135; DB 21; Length 41;  
Best Local Similarity 90.0%; Pred. No. 4.3e-06;  
Matches 27; Conservative 1; Mismatches 2; Indels 0; Gaps 0;  
QY 1 MRLNRLAARPGOPITLLPVRGRKTR 30  
Db 1 MRLNRLAARPGOPITLLPVRGRKTR 30  
RESULT 6  
AAVS8633  
ID AAVS8633 standard; Protein: 1299 AA.  
XX  
XX AAVS8633;  
AC  
XX  
XX 11-APR-2000 (first entry)  
DT  
XX  
DE Protein regulating gene expression PRGE-26.  
XX  
XX Protein regulating gene expression: PRGE-26; human;  
KM cell proliferation; antiproliferative; inflammation;  
KW antiinflammatory; therapy; diagnosis.  
XX  
XX Homo sapiens.  
OS  
XX  
XX  
FH Key Location/Qualifiers  
FT Modified-site 85 /note="O-phosphorylated"  
FT Modified-site 139 /note="O-phosphorylated"  
FT Modified-site 160 /note="O-phosphorylated"  
FT Modified-site 199 /note="O-phosphorylated"  
FT Modified-site 225 /note="O-phosphorylated"  
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FT Modified-site 1001 /note="O-phosphorylated"  
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FT Modified-site 1260 /note="O-phosphorylated"  
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FT Modified-site 43 /note="O-phosphorylated"  
FT Modified-site 136 /note="O-phosphorylated"  
FT Modified-site 268 /note="O-phosphorylated"  
FT Modified-site 168 /note="O-phosphorylated"  
FT Modified-site 91 /note="N-glycosylated"  
FT Modified-site 934 /note="N-glycosylated"

FT Modified-site 991 /note- "N-glycosylated"  
FT Modified-site 1031 /note- "N-glycosylated"  
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FT Modified-site 1235 /note- "N-glycosylated"  
FT Modified-site 1246 /note- "N-glycosylated"  
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XX W09964596-A2.  
XX 16-DEC-1999.  
XX 11-JUN-1999; 99WO-US13281.  
XX 12-JUN-1998; 98US-0089029.  
XX 29-JUL-1998; 98US-0094575.  
XX 14-OCT-1998; 98US-0104624.  
XX (INCY-) INCYTE PHARM INC.  
XX Lal P, Yue H, Tang YT, Hillman JL, Bandman O, Corley NC;  
PI Guejler KJ, Gorgone GA, Baughn MR, Patterson C, Lu DAM;  
XX MPI: 2000-116543/10.  
XX N-PSDB; AA257864.  
XX New human polypeptides that regulate gene expression, for treatment,  
PT prevention and diagnosis of, e.g. cancer -  
XX Claim 1; Page 110-113; 150pp; English.  
XX The present sequence is that of new human protein regulating gene  
CC expression PRGE-26, deduced from incyte clone PITONOT01 (see AA257864)  
CC isolated from pituitary gland cDNA library. PRGE-26 is expressed in  
CC reproductive, nervous and gastrointestinal tissues associated with  
CC cell proliferative and inflammation diseases, disorders or conditions.  
CC It is characterised as a glutamine-rich protein. The invention  
CC provides PRGE polypeptides (see AA258608-38) and polynucleotides (see  
CC AA257839-69), expression vectors, host cells, antibodies, agonists and  
CC antagonists. It also provides methods for diagnosing, treating or  
CC preventing disorders associated with expression of PRGE.  
XX Sequence 1299 AA;  
SO  
Query Match 12.5%; Score 126; DB 21; Length 1299;  
Best Local Similarity 29.1%; Pred. No. 0.0022;  
Matches 44; Conservative 29; Mismatches 62; Indels 16; Gaps 5;  
QY 58 ENRQ---YREYVRLRRFTELEVRGKLEHARAGVLAERKAEALREHGE---LMAWR 110  
DB 754 ektqgeellrrqgeellrrqgeekrreee---larrkgeallrrqgeallrrqge 809  
QY 111 EENRRLQELRIKRLQLEAQAELRQAEVQORAEQA---WVQLKEQVLEKLEAKNFI 168  
DB 810 eeerqgeallrrleerreeertrqgeellrrkgeeeakvaereeaqrlee---nrl 866  
QY 169 TREMLEARIEALDSEKSYMAVTKGQVVR 199  
DB 867 rmeeaarlrheeerkrkelevqrkqlmr 897  
RESULT 7  
AAG07955  
ID AAG07955 standard; Protein; 176 AA.  
XX

AC AAG07955;  
XX 17-OCT-2000 (first entry)  
DE Arabidopsis thaliana protein fragment SEQ ID NO: 5303.  
XX Protein identification; signal transduction pathway; metabolic pathway;  
KW hybridisation assay; genetic mapping; gene expression control; promoter;  
XX termination sequence.  
OS Arabidopsis thaliana.  
XX EPI033405-A2.  
XX 06-SEP-2000.  
XX 25-FEB-2000; 2000EP-0301439.  
XX 25-FEB-1999; 99US-0121825.  
XX 05-MAR-1999; 99US-0123180.  
XX 09-MAR-1999; 99US-0123548.  
XX 23-MAR-1999; 99US-0125788.  
XX 25-MAR-1999; 99US-0126264.  
XX 29-MAR-1999; 99US-0126785.  
XX 01-APR-1999; 99US-0127462.  
XX 06-APR-1999; 99US-0128234.  
XX 08-APR-1999; 99US-0128714.  
XX 16-APR-1999; 99US-0129845.  
XX 19-APR-1999; 99US-0130077.  
XX 21-APR-1999; 99US-0130449.  
XX 23-APR-1999; 99US-0130510.  
XX 28-APR-1999; 99US-0130891.  
XX 30-APR-1999; 99US-0131449.  
XX 30-APR-1999; 99US-0132048.  
XX 04-MAY-1999; 99US-0132407.  
XX 05-MAY-1999; 99US-0132484.  
XX 06-MAY-1999; 99US-0132485.  
XX 07-MAY-1999; 99US-0132486.  
XX 07-MAY-1999; 99US-0132487.  
XX 11-MAY-1999; 99US-0132863.  
XX 14-MAY-1999; 99US-0134256.  
XX 14-MAY-1999; 99US-0134218.  
XX 14-MAY-1999; 99US-0134219.  
XX 14-MAY-1999; 99US-0134221.  
XX 14-MAY-1999; 99US-0134370.  
XX 18-MAY-1999; 99US-0134376.  
XX 19-MAY-1999; 99US-0134941.  
XX 20-MAY-1999; 99US-0135124.  
XX 21-MAY-1999; 99US-0135353.  
XX 24-MAY-1999; 99US-0135629.  
XX 25-MAY-1999; 99US-0136021.  
XX 27-MAY-1999; 99US-0136382.  
XX 28-MAY-1999; 99US-0136782.  
XX 01-JUN-1999; 99US-0137222.  
XX 03-JUN-1999; 99US-0137528.  
XX 04-JUN-1999; 99US-0137529.  
XX 07-JUN-1999; 99US-0137724.  
XX 08-JUN-1999; 99US-0138094.  
XX 10-JUN-1999; 99US-0138540.  
XX 10-JUN-1999; 99US-0138847.  
XX 14-JUN-1999; 99US-0139119.  
XX 16-JUN-1999; 99US-0139452.  
XX 16-JUN-1999; 99US-0139453.  
XX 17-JUN-1999; 99US-0139452.  
XX 18-JUN-1999; 99US-0139454.  
XX 18-JUN-1999; 99US-0139455.  
XX 18-JUN-1999; 99US-0139456.  
XX 18-JUN-1999; 99US-0139457.  
XX 18-JUN-1999; 99US-0139458.  
XX 18-JUN-1999; 99US-0139459.  
XX 18-JUN-1999; 99US-0139460.  
XX 18-JUN-1999; 99US-0139461.  
XX 18-JUN-1999; 99US-0139462.

Query Match	12.5%	Score 125;	DB 21;	Length 176;
Best Local Similarity	29.2%	Pred: No. 0.00023;		
Matches 38;	Conservative 31;	Mismatches 51;	Indels 10;	Gaps 3
OY 56	LTERVROYRETVRALRREFTLEVR---	GKLHEARACVLAERKRAQERIRHODLMMNRE	111	
Db 47	laqrdrlyakkrvyrkyrtheimlrvkqrkdearlllaeraaneerillkaaaakvra		106	
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XX Protein identification: signal transduction pathway; metabolic pathway;  
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Query Match 12.5%; Score 125; DB 21; Length 176;  
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 OY 112 ENRRLOEIRIARLQLEAOELROAEVQORAOEBQAVOLKEQEVYKLOFEAKNFTIRE 171  
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 Db 161 elerkitael 170

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 AC 18-OCT-2000 (first entry)  
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 DE Protein identification; signal transduction pathway; metabolic pathway;  
 KW hybridisation assay; genetic mapping; gene expression control; promoter;  
 KW termination sequence.  
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 OS Arabidopsis thaliana.  
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PR 29-OCT-1999; 99US-0162142.
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Query Match 12.5%; Score 125; DB 21; Length 195;  
Best Local Similarity 29.2%; Pred. No. 0.00026;  
Matches 38; Conservative 31; Mismatches 51; Indels 10; Gaps 3;

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Db 66 laqiriyakkvrykhelemrlvkekqrdearlllaeraaneerlllkaeakvra 125
OY 112 ENRRLOELRLARQLLEQAQDELQAQAEVQAQARADEQAQVQLKQDEVLKLOEAKNFTIRE 171
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OY 172 NLEARIEAL 181
Db 180 elerklteal 189
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RESULT 11
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XX AAG48182;
DT 18-OCT-2000 (first entry)
DE Arabidopsis thaliana protein fragment SEQ ID NO: 60820.
KW Protein identification; signal transduction pathway; metabolic pathway;
KM Hybridisation assay; genetic mapping; gene expression control; promoter;
KW termination sequence.
XX Arabidopsis thaliana.
OS Arabidopsis thaliana.
PN EP1033405-A2.
PD 06-SEP-2000.
XX 25-FEB-2000; 2000EP-0301439.
PF 25-FEB-1999; 99US-0121825.
PR 05-MAR-1999; 99US-0123180.
PR 09-MAR-1999; 99US-0123548.
PR 23-MAR-1999; 99US-0125788.
PR 25-MAR-1999; 99US-0126264.
PR 29-MAR-1999; 99US-0126785.
PR 01-APR-1999; 99US-0127462.
PR 06-APR-1999; 99US-0128234.
PR 08-APR-1999; 99US-0128714.
PR 16-APR-1999; 99US-0129845.
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PR 21-APR-1999; 99US-0130449.
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PR 28-APR-1999; 99US-0131449.
PR 30-APR-1999; 99US-0132048.
PR 30-APR-1999; 99US-0132407.
PR 04-MAY-1999; 99US-0132484.
PR 05-MAY-1999; 99US-0132485.
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AC AAG48203;  
DT 18-OCT-2000 (first entry)  
DE Arabidopsis thaliana protein fragment SEQ ID NO: 60848.  
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XX Protein identification: signal transduction pathway; metabolic pathway;  
KM hybridisation assay; genetic mapping; gene expression control; promoter;  
KM termination sequence.  
XX Arabidopsis thaliana.  
OS  
PN EPI033405-A2.  
PD 06-SEP-2000.  
XX  
XX 25-FEB-2000; 2000EP-0301439.  
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Query Match 12.5% Score 125; DB 21; Length 195;  
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RESULT 13  
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XX  
DT 17-OCT-2000 (first entry)  
XX  
DE Arabidopsis thaliana protein fragment SEQ ID NO: 5301.  
XX  
KW Protein identification; signal transduction pathway; metabolic pathway;  
KW hybridisation assay; genetic mapping; gene expression control; promoter;  
KW termination sequence.  
OS Arabidopsis thaliana.  
PN EP1033405-A2.  
XX  
PD 06-SEP-2000.  
XX  
PE 25-FEB-2000; 2000EP-0301439.  
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Db 251 elerktreal 260

RESULT 14  
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AC AAC48202;  
XX  
DT 18-OCT-2000 (first entry)  
XX  
DE Arabidopsis thaliana protein fragment SEQ ID NO: 60847.  
XX  
KW Protein identification; signal transduction pathway; metabolic pathway;  
KW hydrolisation assay; genetic mapping; gene expression control; promoter;  
KW termination sequence.  
XX  
OS Arabidopsis thaliana.  
XX  
PN EPI033405-A2.  
XX  
PD 06-SEP-2000.  
XX  
PF 25-FEB-2000; 2000EP-0301439.  
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PR	13-SEP-1999;	99US-0153758.
PR	15-SEP-1999;	99US-0154018.
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PR	24-SEP-1999;	99US-0155659.
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PR	14-OCT-1999;	99US-0159329.
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PR	14-OCT-1999;	99US-0159638.
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PR	21-OCT-1999;	99US-0160770.
PR	21-OCT-1999;	99US-0160814.
PR	21-OCT-1999;	99US-0160815.
PR	22-OCT-1999;	99US-0160980.
PR	22-OCT-1999;	99US-0160981.
PR	22-OCT-1999;	99US-0160989.
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PR	26-OCT-1999;	99US-0161361.
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PR	29-OCT-1999;	99US-0162142.

	Query Match	Similarity	Best Local	Matches	Conservative	Mismatches	Indels	Gaps
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		29.2%;	Pred. No. 0.00039;					
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Oy	56	LTERTYROYRETFVRLARREFTLEVR----	GKLTLEARAGVLAERKAOEALIREHOLMANNEE	111				
Db	137	laqltrdyakkrvyrkeyihneimlreykqkdearliraaneerlllkaaeakvra		196				
Oy	112	ENRRLOETRIARIALQLEAQAOLRQAEEVQAQRAQECQAWOLKEQEVYLKLOEERKKNFTIE		171				
Db	197	eerktradeefrftlllkerereklemvkmmgqkree-----kikerekl-l-iregsiwiqeq		250				
Oy	172	NLEARIEEAL		181				

Db	251	elerk	teal	260
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ID	AAAG48181	standard;	Protein;	280 AA.
AC	AAAG48181;			
XX				
DT	18-OCT-2000	(first entry)		
XX				
DE	Arabidopsis thaliana	protein fragment SEQ ID NO: 60819.		
XX				
KW	Protein identification assay;	signal transduction pathway; metabolic pathway;		
KW	hybridisation assay;	genetic mapping; gene expression control; promoter;		
KW	termination sequence.			
XX				
OS	Arabidopsis thaliana.			
XX				
PN	EP1033405-A2.			
XX				
PD	06-SEP-2000.			
XX				
PF	25-FEB-2000;	2000EP-0301439.		
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PR	25-FEB-1999;	99US-0121825.		
PR	05-MAR-1999;	99US-0123180.		
PR	09-MAR-1999;	99US-0123548.		
PR	23-MAR-1999;	99US-0125788.		
PR	25-MAR-1999;	99US-0126264.		
PR	29-MAR-1999;	99US-0126785.		
PR	01-APR-1999;	99US-0127462.		
PR	06-APR-1999;	99US-0128234.		
PR	08-APR-1999;	99US-0128714.		
PR	16-APR-1999;	99US-0129845.		
PR	19-APR-1999;	99US-0130077.		
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PR	05-MAY-1999;	99US-0132485.		
PR	06-MAY-1999;	99US-0132486.		
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PR	14-MAY-1999;	99US-0134218.		
PR	14-MAY-1999;	99US-0134219.		
PR	14-MAY-1999;	99US-0134221.		
PR	14-MAY-1999;	99US-0134370.		
PR	18-MAY-1999;	99US-0134768.		
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PR	08-JUN-1999;	99US-0138094.		
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PR 06-JUL-1999; 99US-0142390.  
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PR 22-JUL-1999; 99US-0145145.  
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PR 26-JUL-1999; 99US-0145276.  
PR 27-JUL-1999; 99US-0145913.  
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PR 27-JUL-1999; 99US-0145919.  
PR 28-JUL-1999; 99US-0145951.  
PR 28-JUL-1999; 99US-0146386.  
PR 02-AUG-1999; 99US-0146388.  
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PR 03-AUG-1999; 99US-0147038.  
PR 04-AUG-1999; 99US-0147204.  
PR 04-AUG-1999; 99US-0147302.  
PR 05-AUG-1999; 99US-0147392.  
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PR 06-AUG-1999; 99US-0147303.  
PR 06-AUG-1999; 99US-0147416.  
PR 09-AUG-1999; 99US-0147493.  
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PR 27-AUG-1999; 99US-0151065.  
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Query Match 12.5%; Score 125; DB 21; Length 280;

Best Local Similarity 29.2%; Pred. No. 0.00041; Mismatches 51; Indels 10; Gaps 3;

Matches 38; Conservative 31; Mismatches 51; Indels 10; Gaps 3;

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QY 112 ENRILCELRIRALQLEAOERQAEVQAOERQAEQAWQLEKQEVLYKQAEAKNFIRE 171  
DB 211 eerrlaadeefrqltkeraeklemwkmungkree-----kikerekl-lregsslwieqk 264

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Search completed: June 26, 2001, 16:36:27  
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